

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10|799,749
Source: IFWO
Date Processed by STIC: 10/14/2005

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IFWO

RAW SEQUENCE LISTING

DATE: 10/14/2005

PATENT APPLICATION: US/10/799,749

TIME: 09:40:58

Input Set : N:\CrF3\RULE60\10799749.raw

Output Set: N:\CRF4\10142005\J799749.raw

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1 <110> APPLICANT: Kreiswirth, Barry N
2   Nadich, Steven M
3 <120> TITLE OF INVENTION: System and Method for Tracking and Controlling Infections
4 <130> FILE REFERENCE: 19124.0002
5 <140> CURRENT APPLICATION NUMBER: US/10/799,749
6 <141> CURRENT FILING DATE: 2004-03-15
W--> 7 <140> CURRENT APPLICATION NUMBER: PRIOR APPLICATION NUMBER: US/10/073,256
W--> 8 <141> CURRENT FILING DATE: PRIOR FILING DATE: 2002-02-13
9 <160> NUMBER OF SEQ ID NOS: 80
10 <170> SOFTWARE: PatentIn version 3.1
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13 <211> LENGTH: 24
14 <212> TYPE: DNA
15 <213> ORGANISM: Staphylococcus aureus
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29 <213> ORGANISM: Staphylococcus aureus
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34 <211> LENGTH: 24
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36 <213> ORGANISM: Staphylococcus aureus
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45   aaagaagaca acaacaagcc tggt                                24
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48 <211> LENGTH: 24
49 <212> TYPE: DNA

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50 <213> ORGANISM: Staphylococcus aureus
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54 <210> SEQ ID NO: 7
55 <211> LENGTH: 24
56 <212> TYPE: DNA
57 <213> ORGANISM: Staphylococcus aureus
58 <400> SEQUENCE: 7
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62 <211> LENGTH: 24
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64 <213> ORGANISM: Staphylococcus aureus
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66     aaagaagacg gcaacaaacc tggc                                24
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69 <211> LENGTH: 24
70 <212> TYPE: DNA
71 <213> ORGANISM: Staphylococcus aureus
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73     aaagaagacg gcaacaagcc tggc                                24
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76 <211> LENGTH: 240
77 <212> TYPE: DNA
78 <213> ORGANISM: Staphylococcus aureus
79 <400> SEQUENCE: 10
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81     aacaagcctg gtaaagaaga caacaacaaa cctggtaaag aagacggcaa caagcctggc    120
82     aaagaagaca acaacaaacc tggcaaagaa gacggcaaca agcctggtaa agaagacaac    180
83     aacaagcctg gtaaagaaga cggaacaag cctggtaaag aagacggcaa caaacctggc    240
85 <210> SEQ ID NO: 11
86 <211> LENGTH: 10
87 <212> TYPE: DNA
88 <213> ORGANISM: Staphylococcus aureus
89 <400> SEQUENCE: 11
90     attcatagat                                10
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93 <211> LENGTH: 10
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95 <213> ORGANISM: Staphylococcus aureus
96 <400> SEQUENCE: 12
97     cgtactatcc                                10
99 <210> SEQ ID NO: 13
100 <211> LENGTH: 10
101 <212> TYPE: DNA
102 <213> ORGANISM: Staphylococcus aureus
103 <400> SEQUENCE: 13
104     attcgttata                                10
106 <210> SEQ ID NO: 14

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114 <211> LENGTH: 10
115 <212> TYPE: DNA
116 <213> ORGANISM: Staphylococcus aureus
117 <400> SEQUENCE: 15
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122 <212> TYPE: DNA
123 <213> ORGANISM: Staphylococcus aureus
124 <400> SEQUENCE: 16
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127 <210> SEQ ID NO: 17
128 <211> LENGTH: 12
129 <212> TYPE: DNA
130 <213> ORGANISM: Staphylococcus aureus
131 <400> SEQUENCE: 17
132     aattcgcccta gg                                12
134 <210> SEQ ID NO: 18
135 <211> LENGTH: 12
136 <212> TYPE: DNA
137 <213> ORGANISM: Staphylococcus aureus
138 <400> SEQUENCE: 18
139     aattccccta gg                                  12
141 <210> SEQ ID NO: 19
142 <211> LENGTH: 8
143 <212> TYPE: DNA
144 <213> ORGANISM: Staphylococcus aureus
145 <400> SEQUENCE: 19
146     taggccgt                                       8
148 <210> SEQ ID NO: 20
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150 <212> TYPE: DNA
151 <213> ORGANISM: Staphylococcus aureus
152 <400> SEQUENCE: 20
153     ttaaaggcct ga                                  12
155 <210> SEQ ID NO: 21
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157 <212> TYPE: DNA
158 <213> ORGANISM: Staphylococcus aureus
159 <400> SEQUENCE: 21
160     gggtccaata at                                  12
162 <210> SEQ ID NO: 22
163 <211> LENGTH: 8

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166 <400> SEQUENCE: 22
167      ggттаacc
169 <210> SEQ ID NO: 23
170 <211> LENGTH: 1070
171 <212> TYPE: DNA
172 <213> ORGANISM: Enterococcus faecalis
173 <400> SEQUENCE: 23
174      tttttcttgg caattttggt cgtattatcc gcttttttga ctgttcctga cgattcttga      60
175      tttgtctgta tctgtttagt tgcttggttt tctgctactg attcctttgt ttgactagcc      120
176      ttggcagagg gctttgaatt actttgagcg tcattaggat cttgattaga ctctaccgcg      180
177      taaatgacag aattctggcc tttgctttgg ctactttcgt ttacagtgct tggggtgcta      240
178      ctctcaactg tattgttggt tgcgctgggt gtacttgaag cactactttc gctgggtacta      300
179      cttgttttac tggttgtact tgggtgtgtg ctttcatttg tattgcttgt ttcacttgtc      360
180      gtacttgagg tactactttc gctggtacta ctggtttcgc tggttgtgct tggcggtgtg      420
181      ctttcactcg tactactgct ctcaactgtc gtgcttggcg tgctgctttc gcttgattta      480
182      ctggtttcac ttgtcgtgct tgagggtgctg ctttcgctgg tactactgct ctcaactgtc      540
183      gtgcttggcg tgctgctttc gctggtacta ctgctttcac ttgtcgtgct tgagggtacta      600
184      ctttcgcttg tattactggt ttcgctagtt gtacttggtg tggtgctttc atttgatttg      660
185      cttgtttcac ttgtcgtgct tgagggtgctg ctttcgcttg tattactggt tttactgggt      720
186      gtacttggtg tattgctttc atttgatttg cttgtttcac ttgttgact tgagggtgctg      780
187      ctttcgcttg tattactggt ttcactgggt gtgcttgagg tgctgctttc gcttggtgta      840
188      ctagttgtct ctgttgattt ttcaactaac agaagtaacg ccgtttttat gggttggtgt      900
189      aattgattaa tacgcttttg tgcatctgca ggcgttttaa agccaccaag tggtggctct      960
190      aataattctt catctgacca agcaagcagt tggtgtaact gcttagagct tccttcgcca      1020
191      gttgttgat ctattaaggc ttcttgcatg gcttgccaag agtctttggt      1070
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194 <211> LENGTH: 45
195 <212> TYPE: DNA
196 <213> ORGANISM: Enterococcus faecalis
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200 <210> SEQ ID NO: 25
201 <211> LENGTH: 45
202 <212> TYPE: DNA
203 <213> ORGANISM: Enterococcus faecalis
204 <400> SEQUENCE: 25
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207 <210> SEQ ID NO: 26
208 <211> LENGTH: 45
209 <212> TYPE: DNA
210 <213> ORGANISM: Enterococcus faecalis
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214 <210> SEQ ID NO: 27
215 <211> LENGTH: 45
216 <212> TYPE: DNA
217 <213> ORGANISM: Enterococcus faecalis

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Input Set : N:\Crf3\RULE60\10799749.raw

Output Set: N:\CRF4\10142005\J799749.raw

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219      gtacttgagg tactactttc gctggtacta ctggtttcgc tggtt      45
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222 <211> LENGTH: 45
223 <212> TYPE: DNA
224 <213> ORGANISM: Enterococcus faecalis
225 <400> SEQUENCE: 28
226      gtgcttgagg tggtgctttc actcggtacta ctgctctcac ttgtc      45
228 <210> SEQ ID NO: 29
229 <211> LENGTH: 45
230 <212> TYPE: DNA
231 <213> ORGANISM: Enterococcus faecalis
232 <400> SEQUENCE: 29
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235 <210> SEQ ID NO: 30
236 <211> LENGTH: 45
237 <212> TYPE: DNA
238 <213> ORGANISM: Enterococcus faecalis
239 <400> SEQUENCE: 30
240      gtgcttgagg tgctgctttc gctggtacta ctgctctcac ttgtc      45
242 <210> SEQ ID NO: 31
243 <211> LENGTH: 45
244 <212> TYPE: DNA
245 <213> ORGANISM: Enterococcus faecalis
246 <400> SEQUENCE: 31
247      gtgcttgagg tgctgctttc gctggtacta ctgctttcac ttgtc      45
249 <210> SEQ ID NO: 32
250 <211> LENGTH: 45
251 <212> TYPE: DNA
252 <213> ORGANISM: Enterococcus faecalis
253 <400> SEQUENCE: 32
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256 <210> SEQ ID NO: 33
257 <211> LENGTH: 45
258 <212> TYPE: DNA
259 <213> ORGANISM: Enterococcus faecalis
260 <400> SEQUENCE: 33
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263 <210> SEQ ID NO: 34
264 <211> LENGTH: 45
265 <212> TYPE: DNA
266 <213> ORGANISM: Enterococcus faecalis
267 <400> SEQUENCE: 34
268      gtacttggtg tattgctttc atttggtattg cttggtttcac ttgtt      45
270 <210> SEQ ID NO: 35
271 <211> LENGTH: 45
272 <212> TYPE: DNA
273 <213> ORGANISM: Enterococcus faecalis
274 <400> SEQUENCE: 35

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/799,749

DATE: 10/14/2005
TIME: 09:40:59

Input Set : N:\Crf3\RULE60\10799749.raw
Output Set: N:\CRF4\10142005\J799749.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 3

VERIFICATION SUMMARY

DATE: 10/14/2005

PATENT APPLICATION: US/10/799,749

TIME: 09:40:59

Input Set : N:\Crf3\RULE60\10799749.raw

Output Set: N:\CRF4\10142005\J799749.raw

L:7 M:280 W: Numeric Identifier already exists, <140> found multiple times
L:7 M:281 W: Numeric Fields not Ordered, <140> not ordered!.
L:7 M:270 C: Current Application Number differs, Replaced Current Application Number
L:8 M:280 W: Numeric Identifier already exists, <141> found multiple times
L:8 M:256 W: Invalid Numeric Header Field, Wrong Current FILING DATE:YYYY-MM-DD